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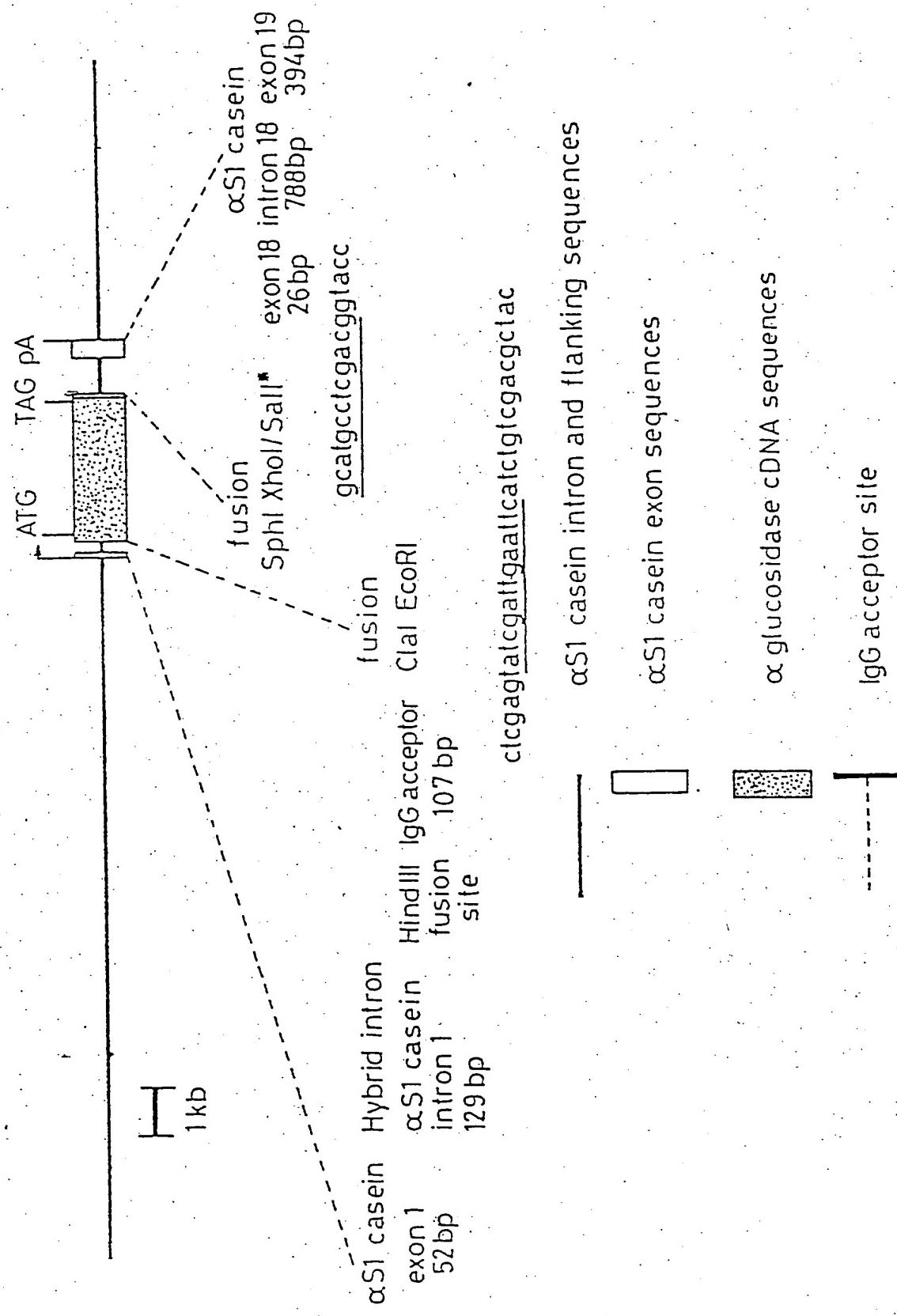
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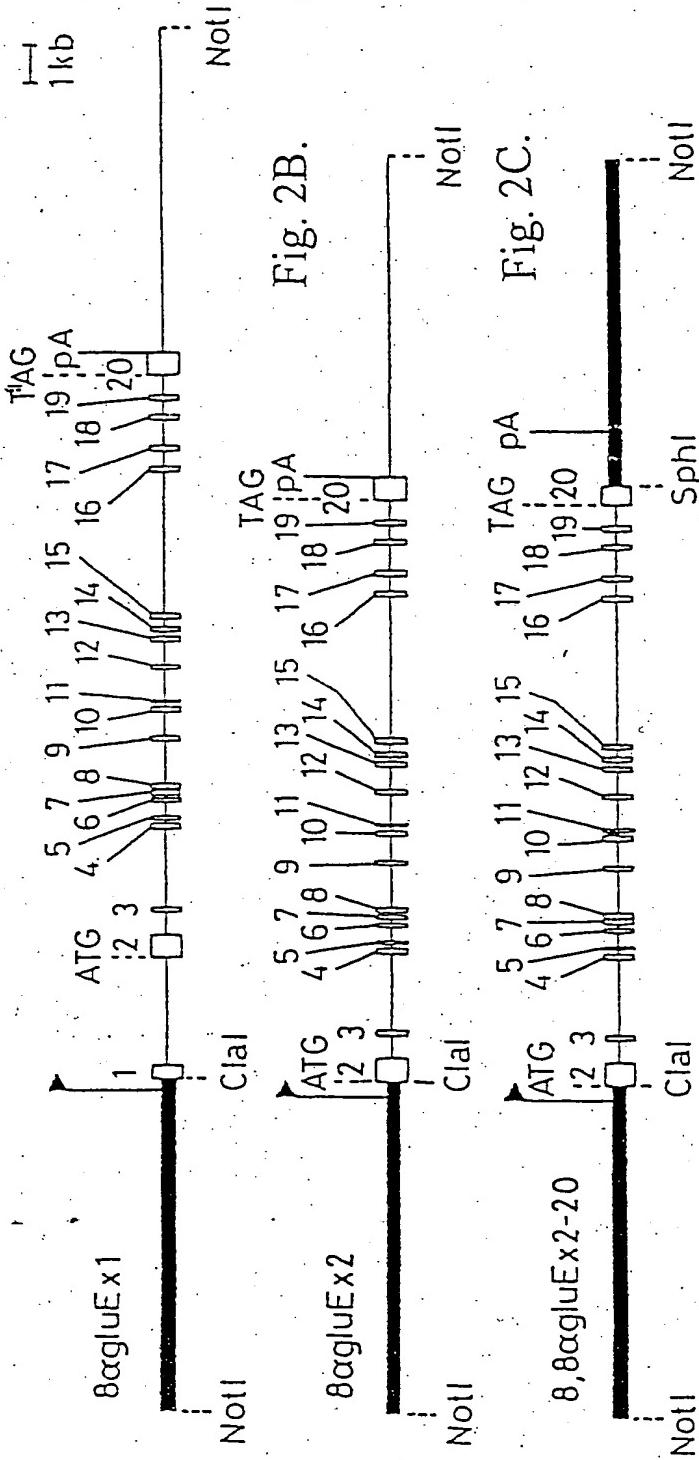
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Fig. 1.



α -glucosidase constructs

Fig. 2.A



Transcription Initiation site.

■ αs_1 casein sequence, promoter or 3' untranslated region.

- 2 3 The boxes represent the exons in the α -glucosidase sequence, the thin line represents the intron sequences.
 □ - The numbers above the boxes are the exon numbers
 PA = polyadenylation signal.
 ATG = translation initiation site.
 TAG = translation stop codon

Fig. 2.C.

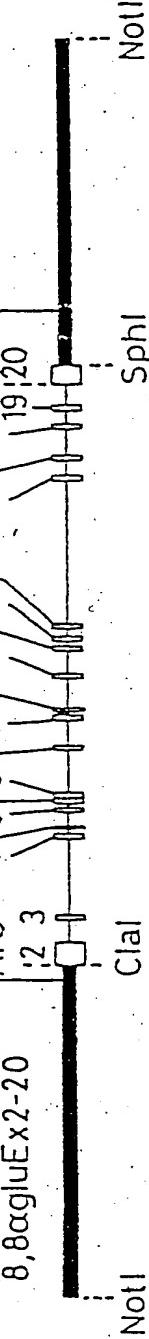


Fig. 2.B.

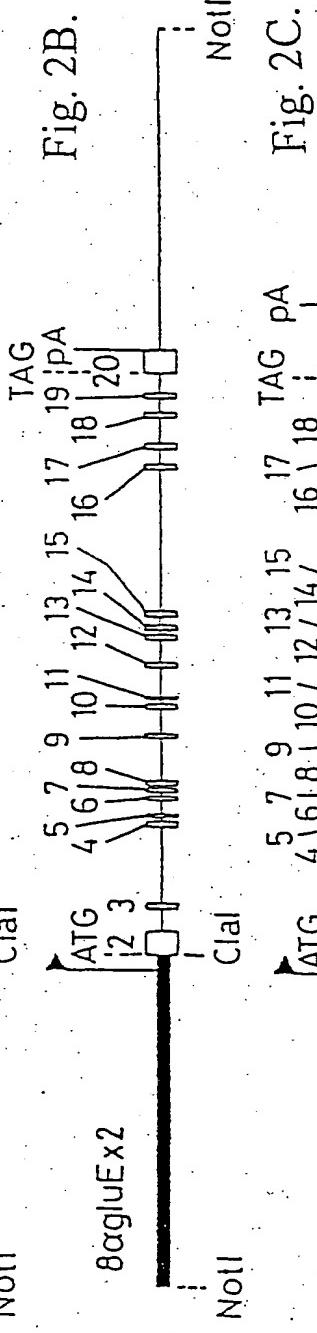


Fig. 3A.

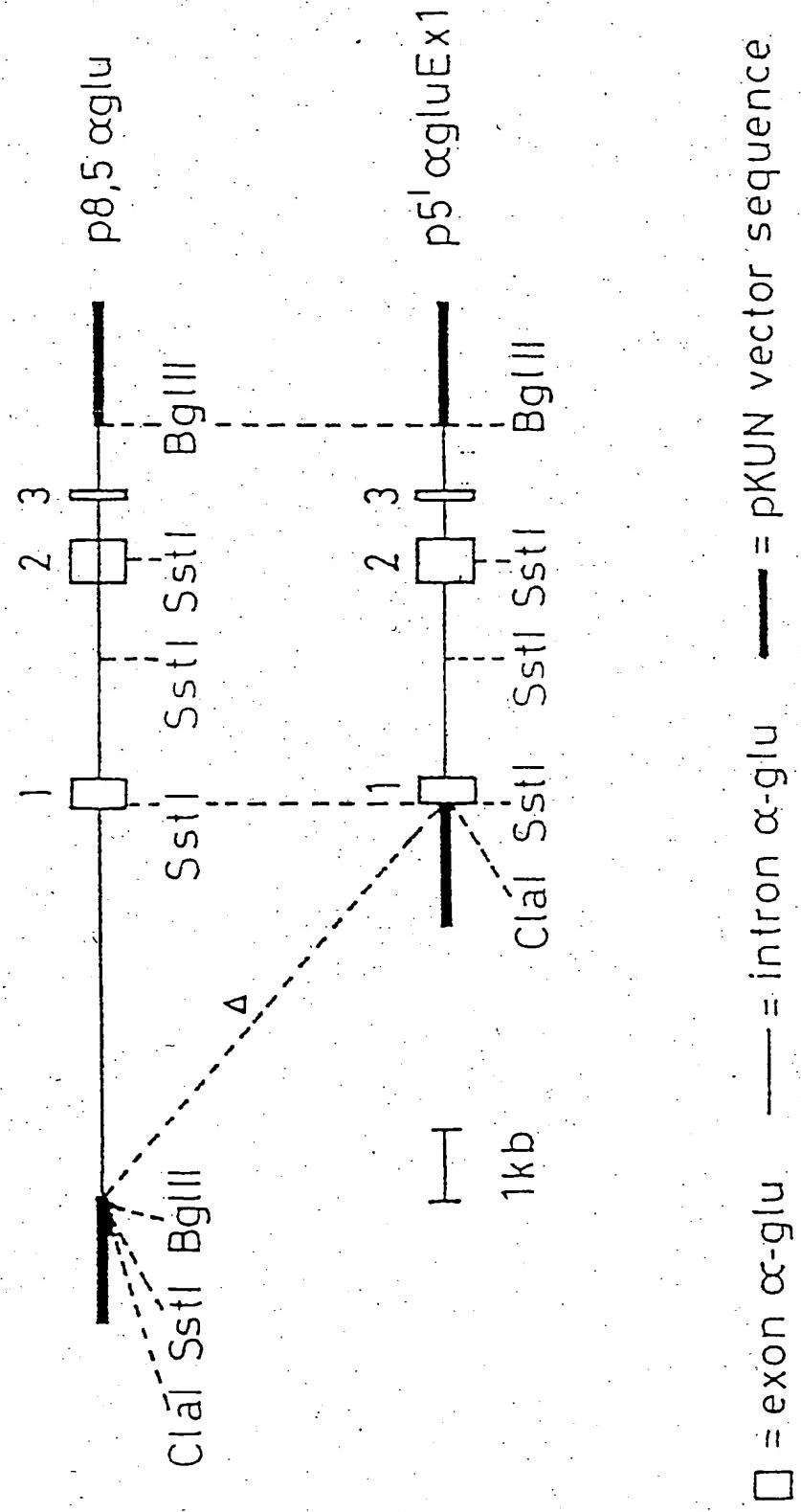


Fig. 3B.

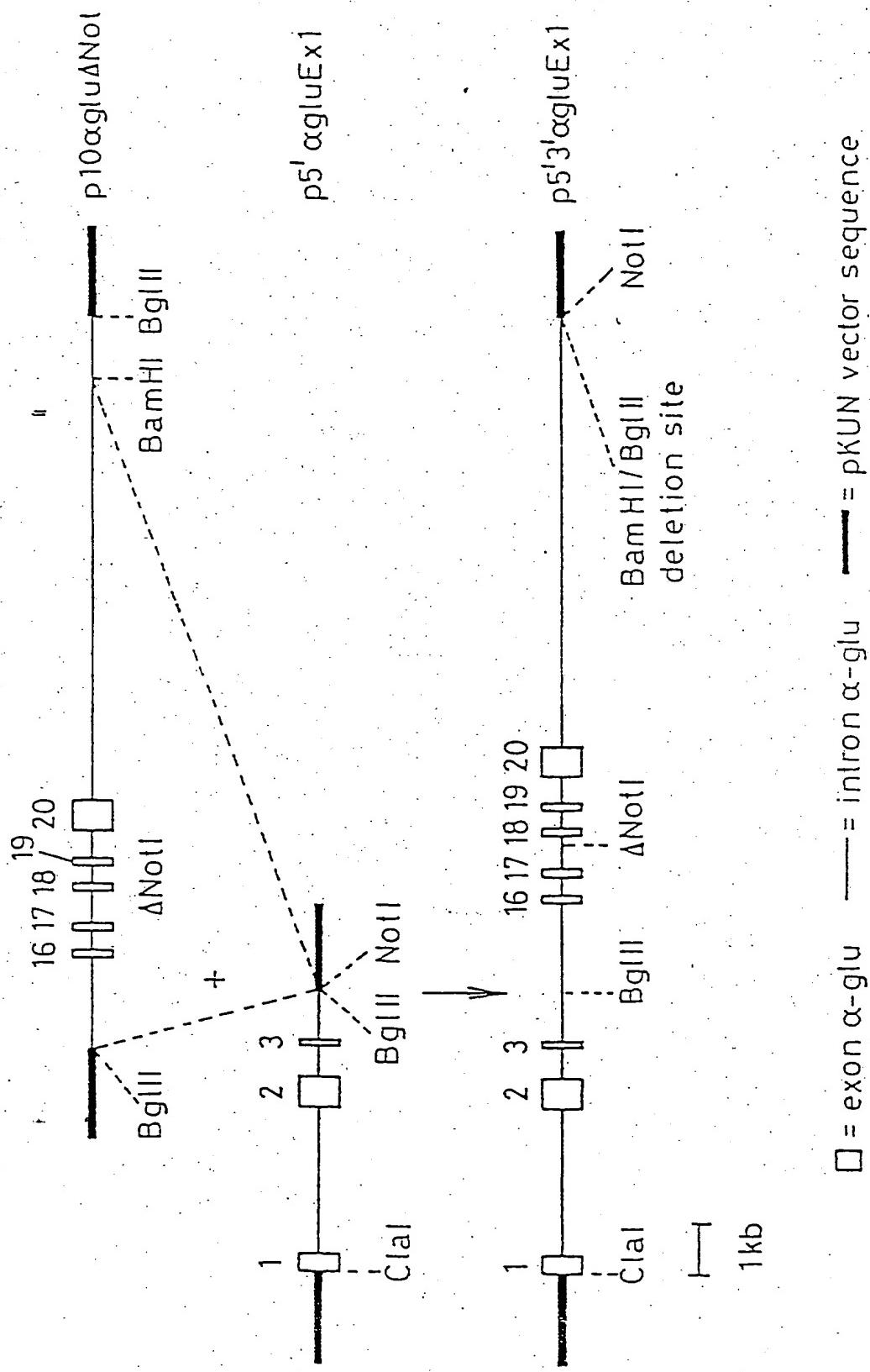


Fig. 3.C.

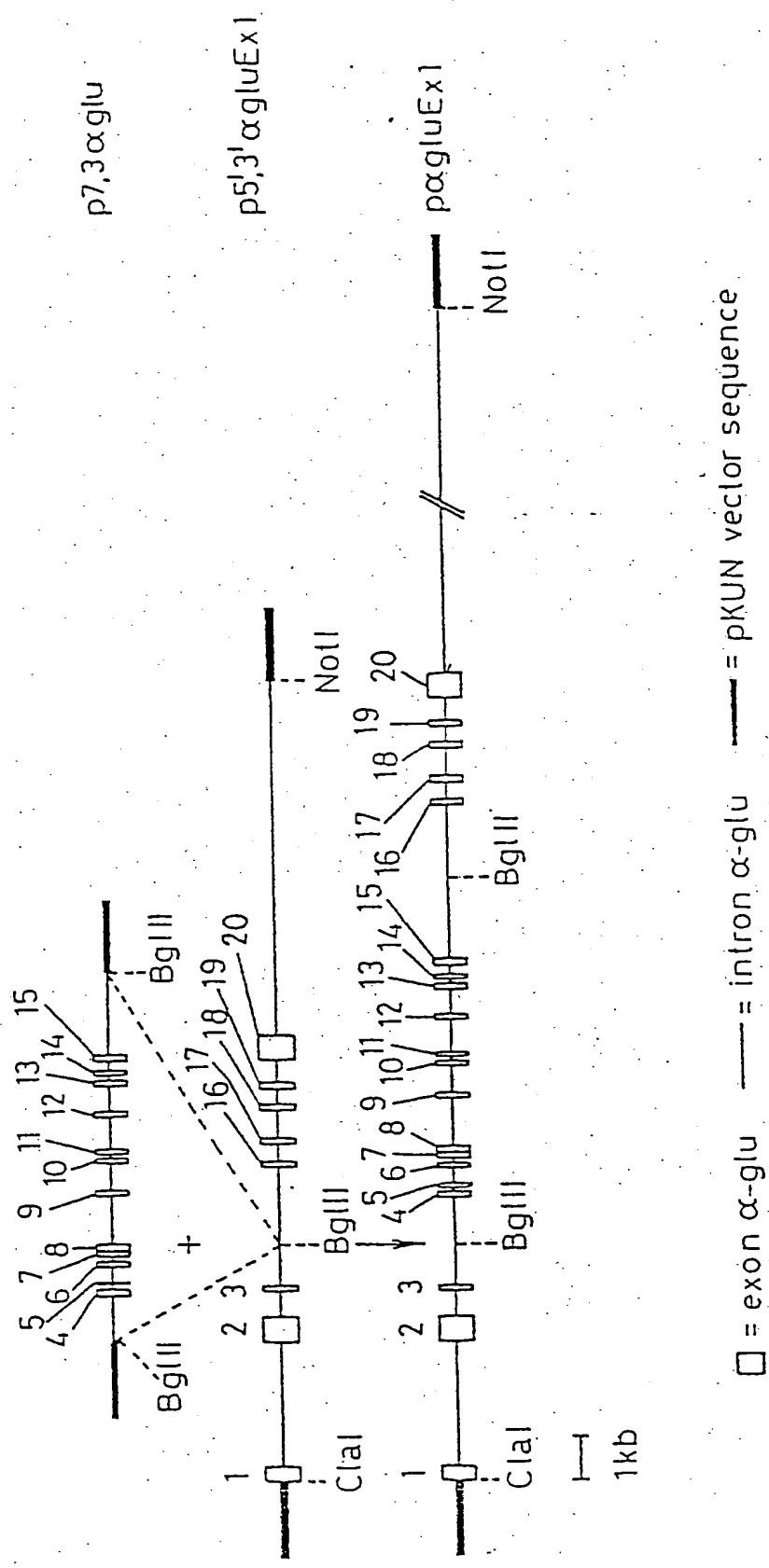


Fig. 4. A.

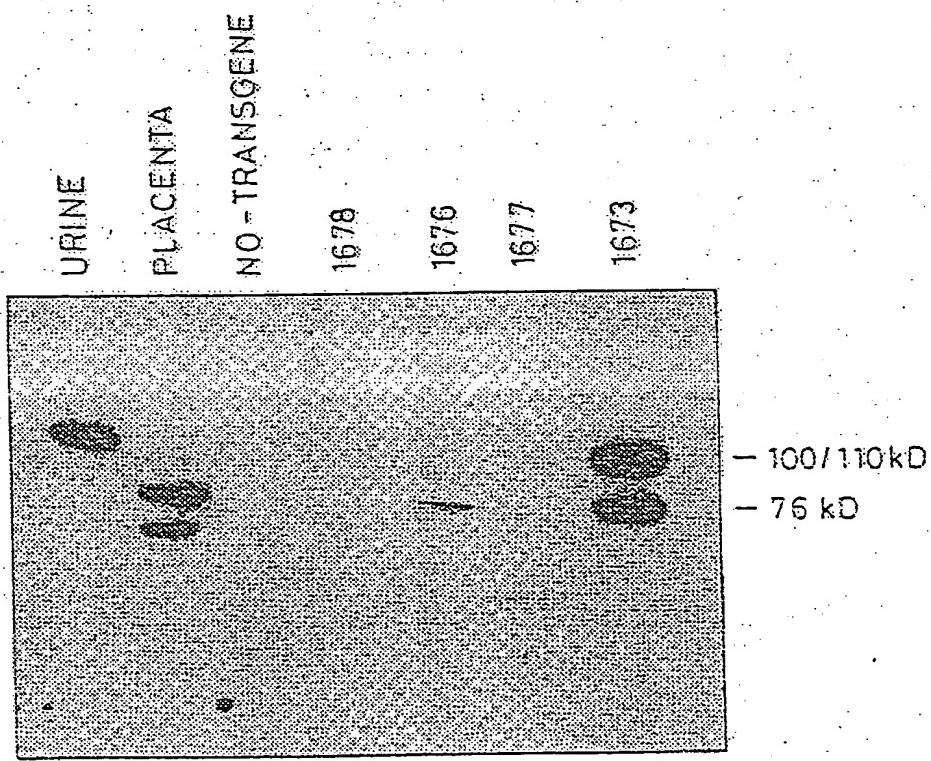


Fig. 4. B.

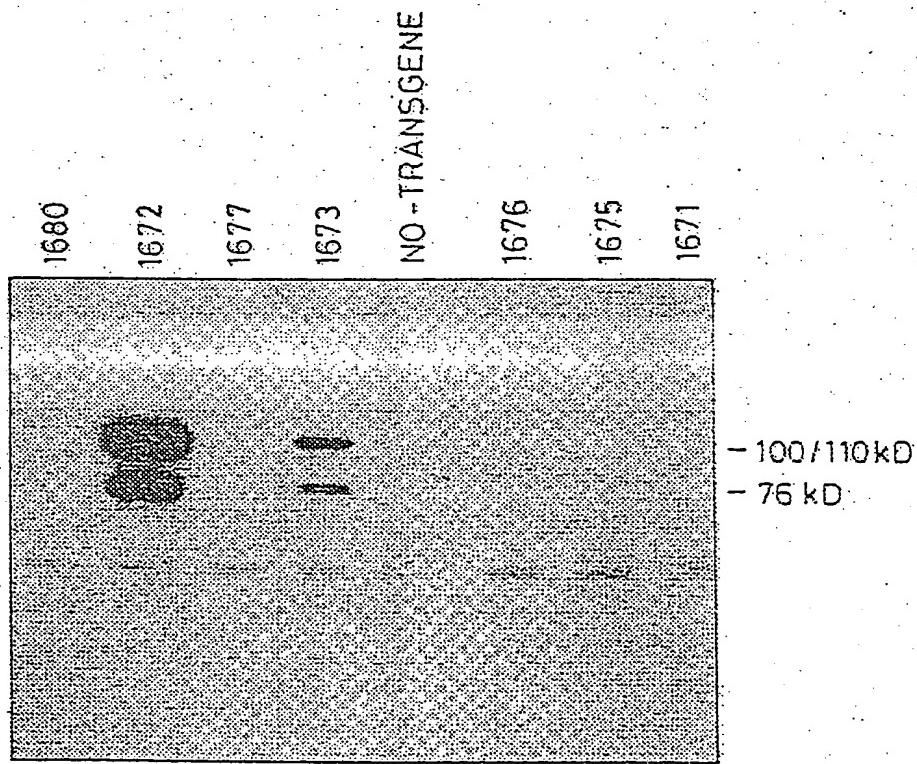


Fig. 5.

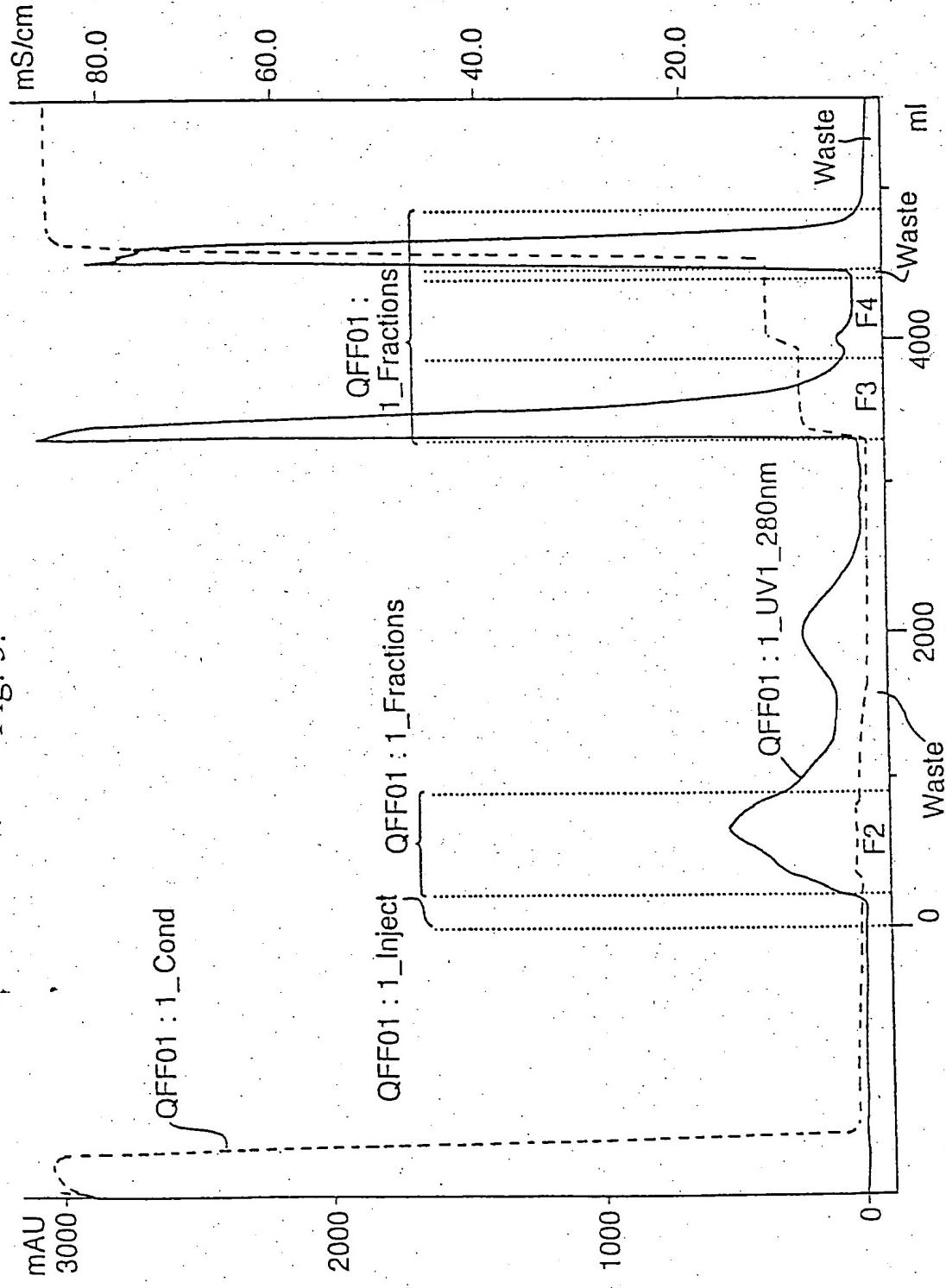


Fig. 6.

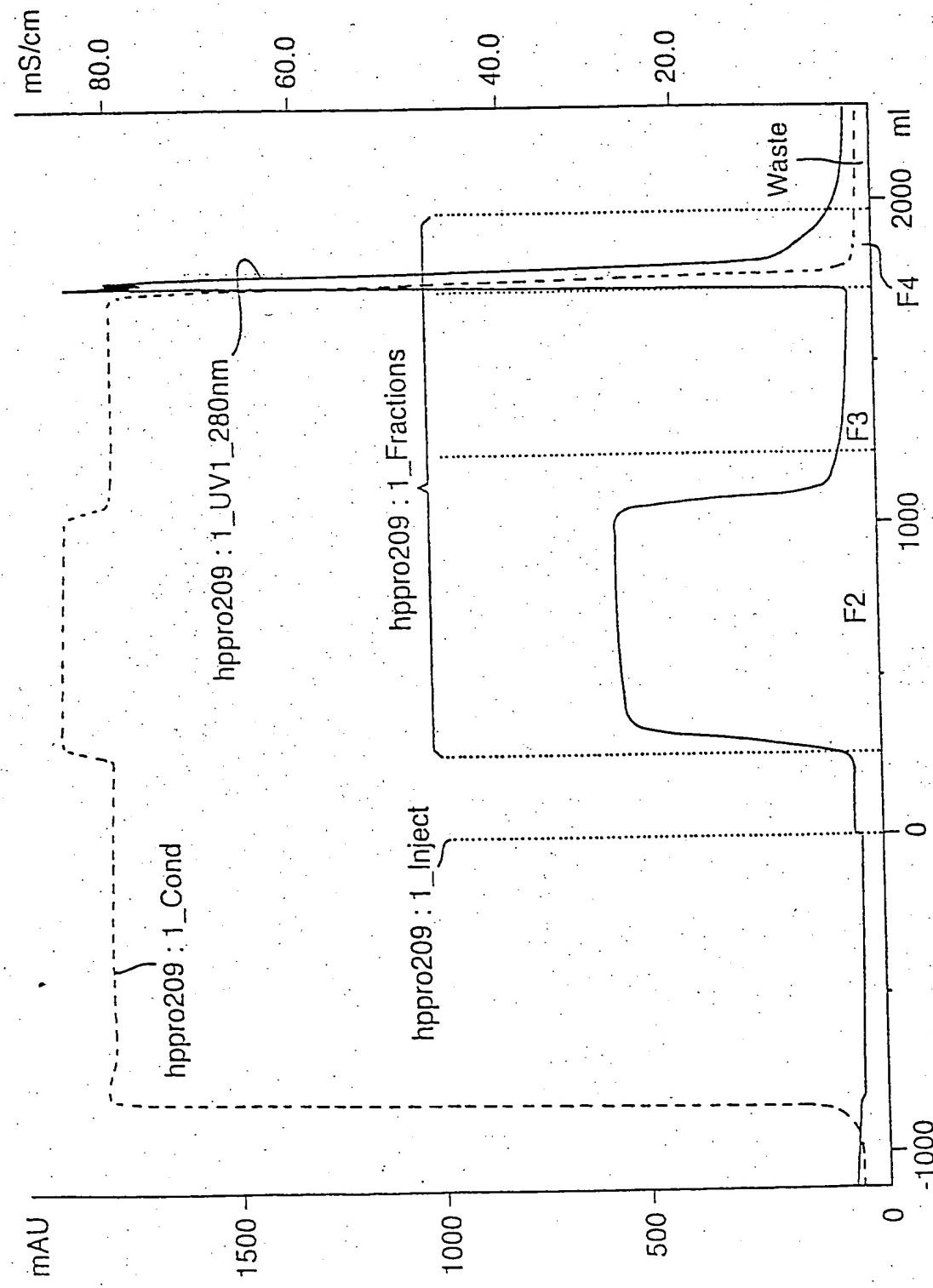


Fig. 7.

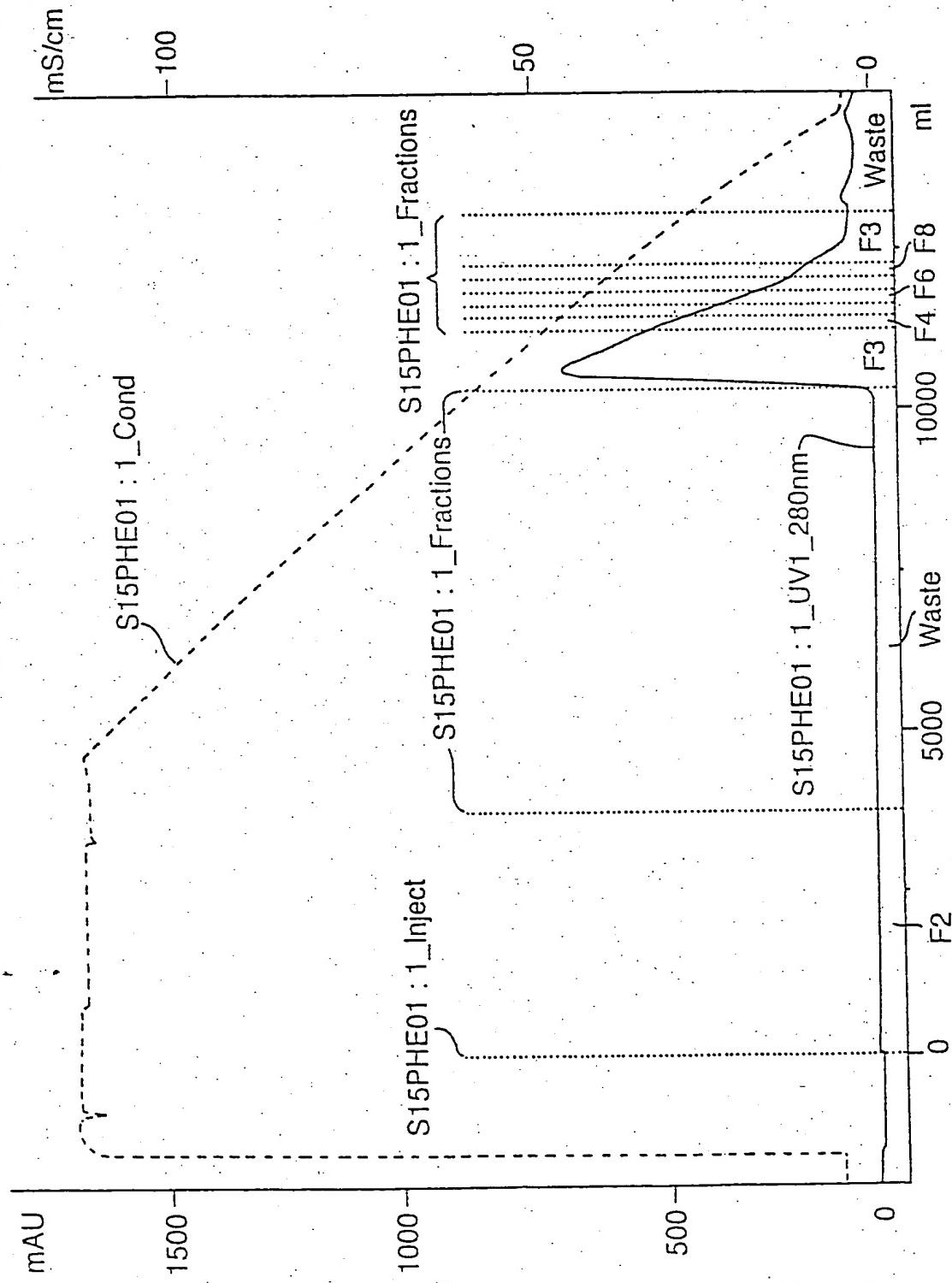


Fig. 8.

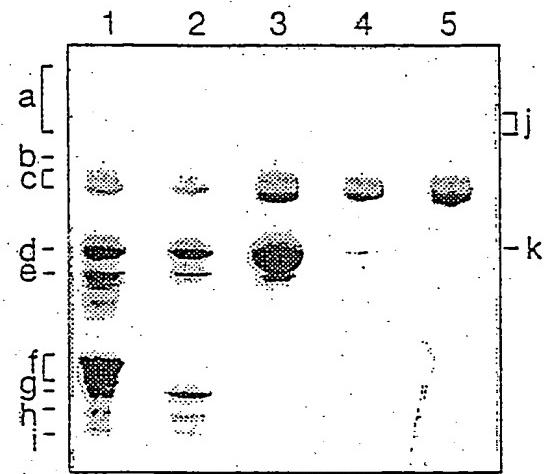


Fig. 9.

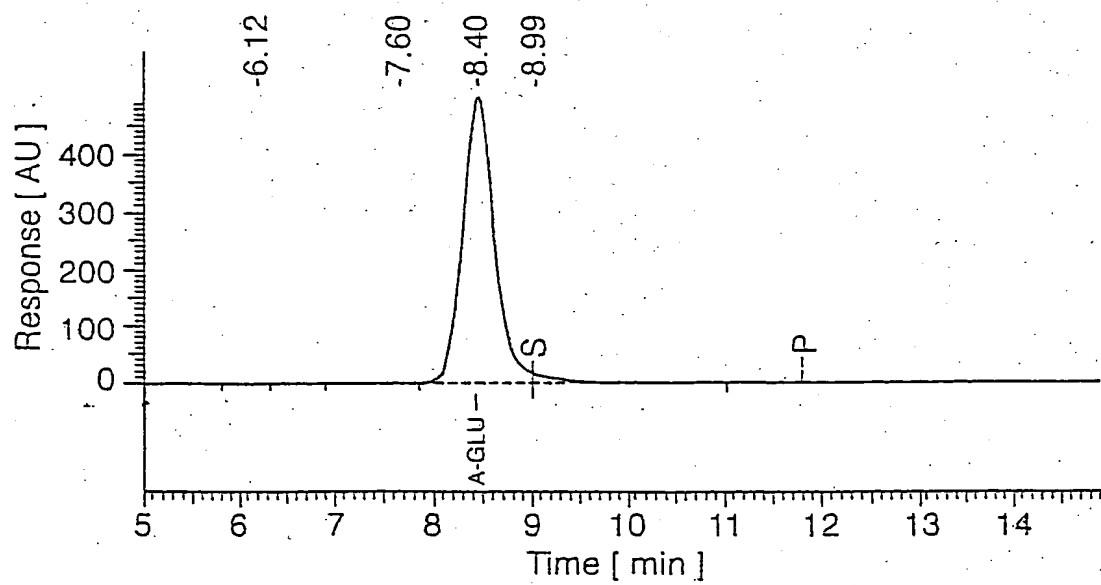


Fig. 10.

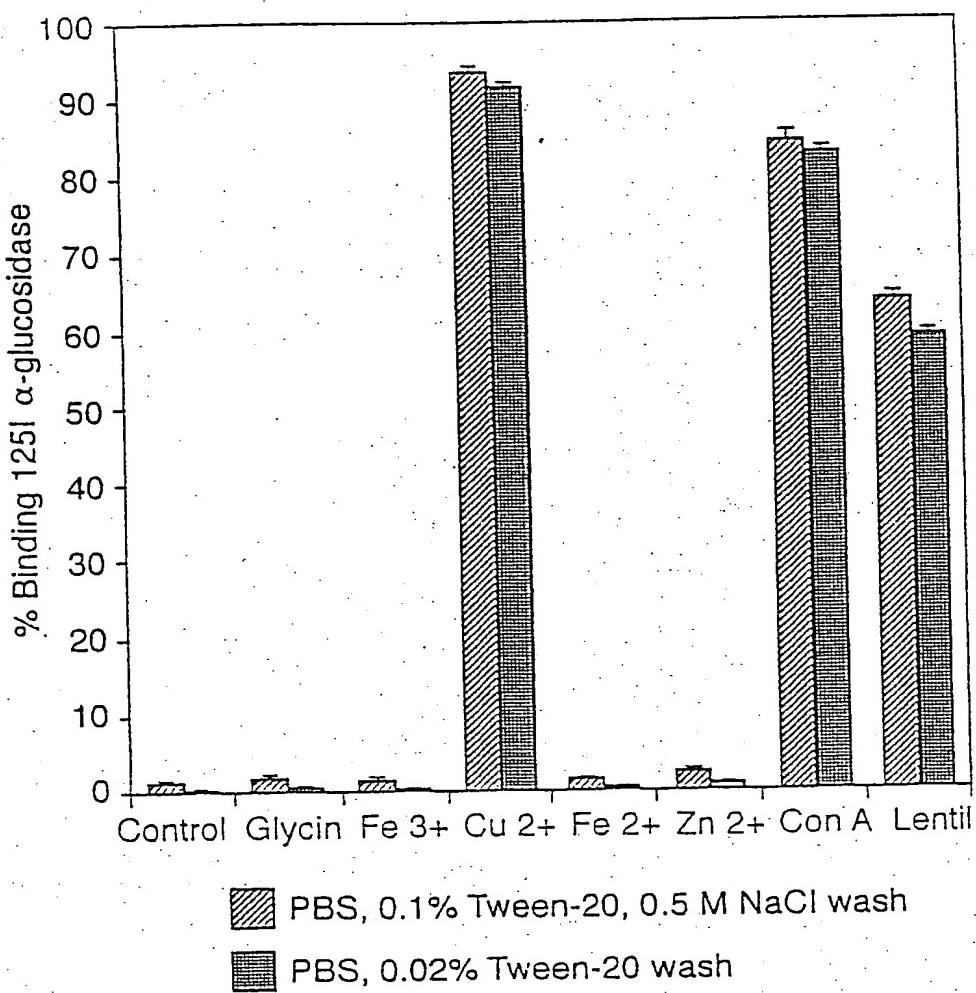


Fig. 11. A.

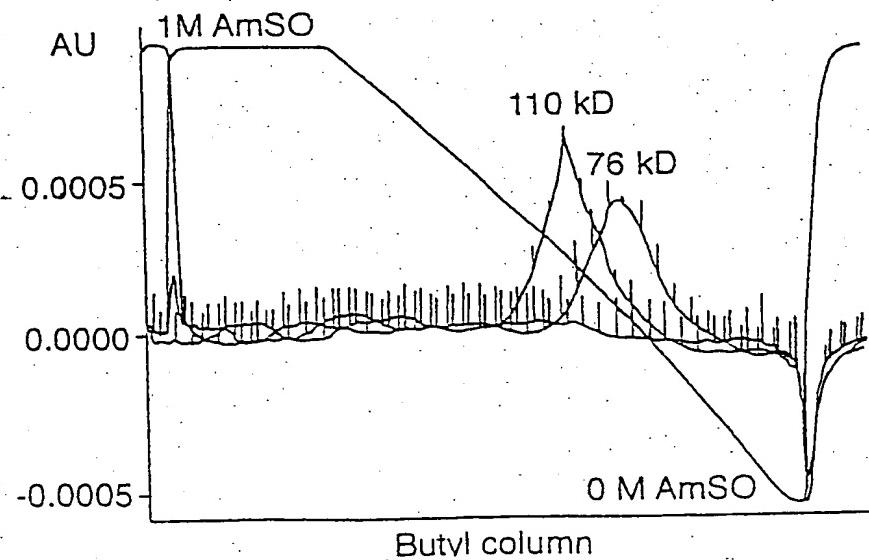


Fig. 11. B.

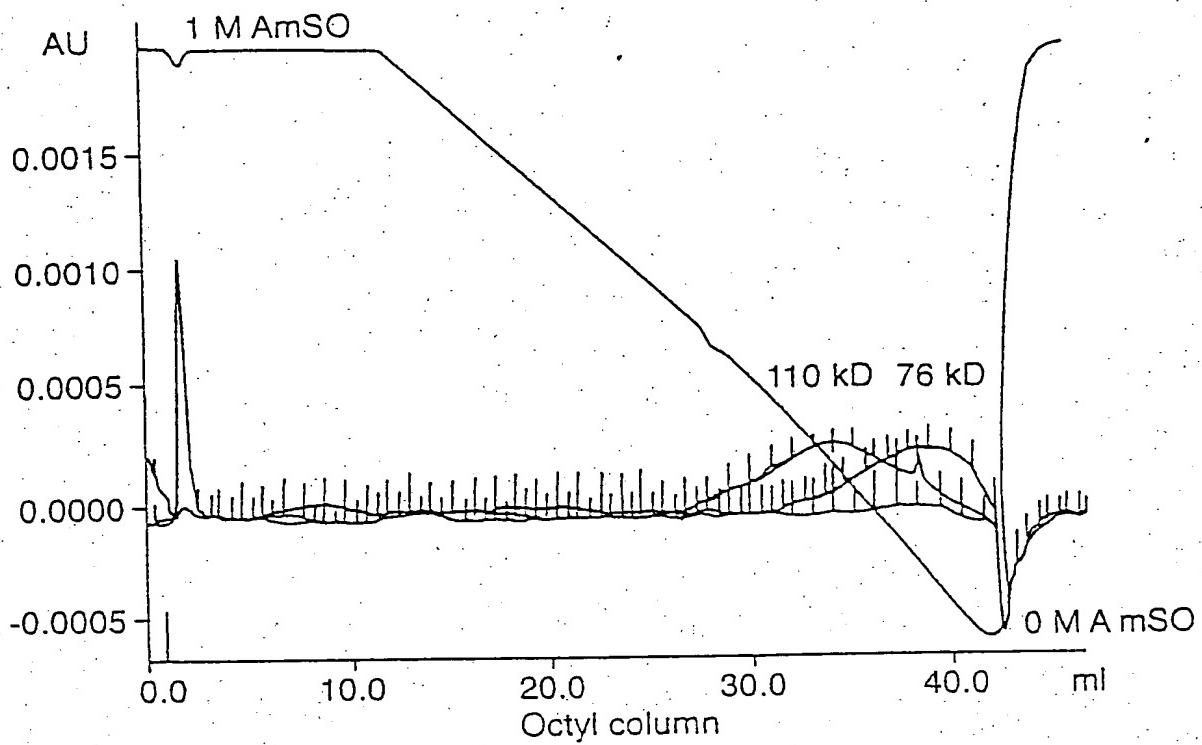


Fig. 11. C.

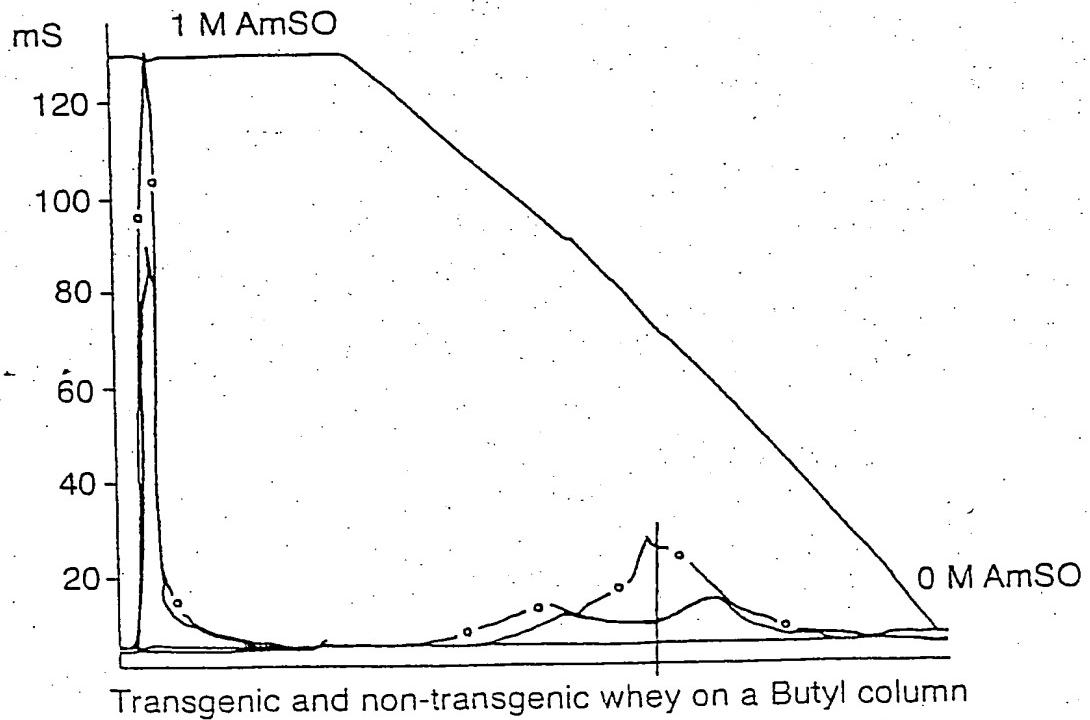


Fig. 11. D.

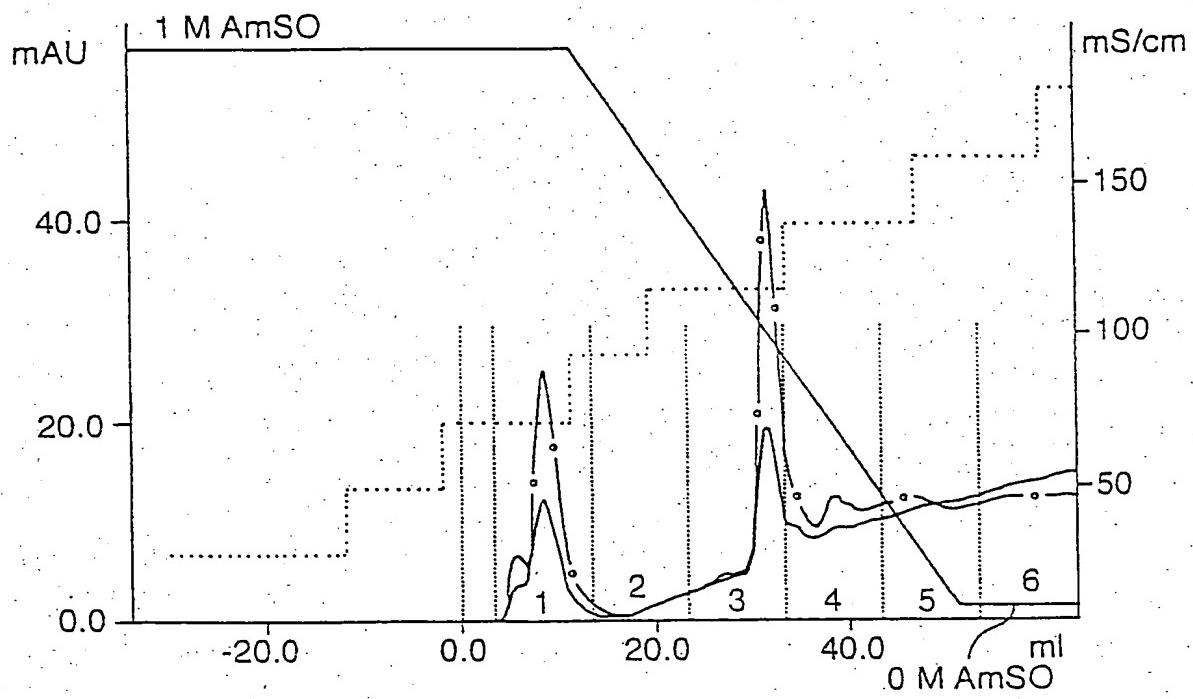


Fig. 12.

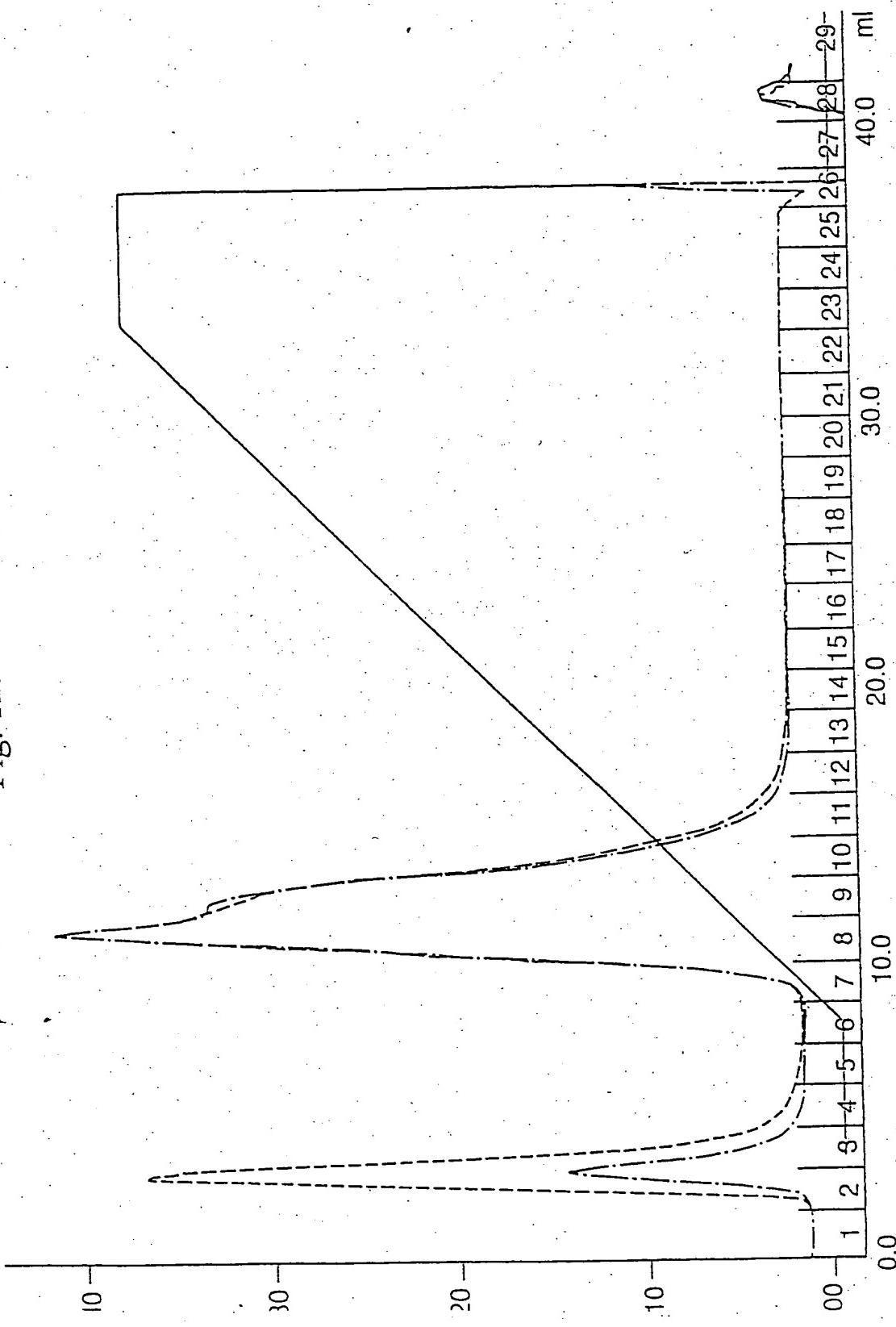
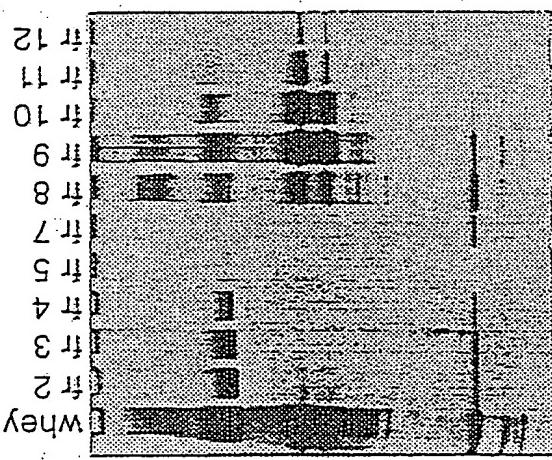


Fig. 13. A.

transgenic whey

(1-40)



α-gli =

Fig. 13. B.

non-transgenic whey

(1-40)

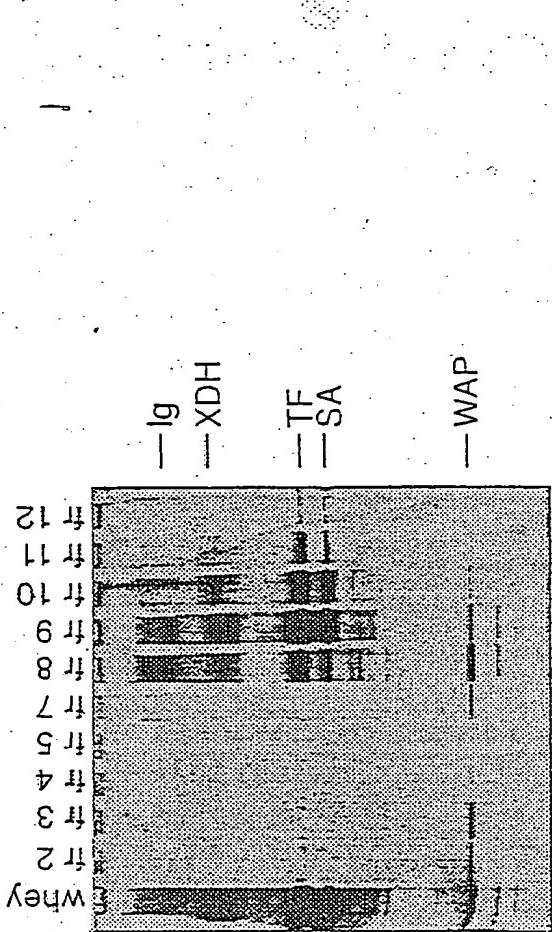


Fig. 14.

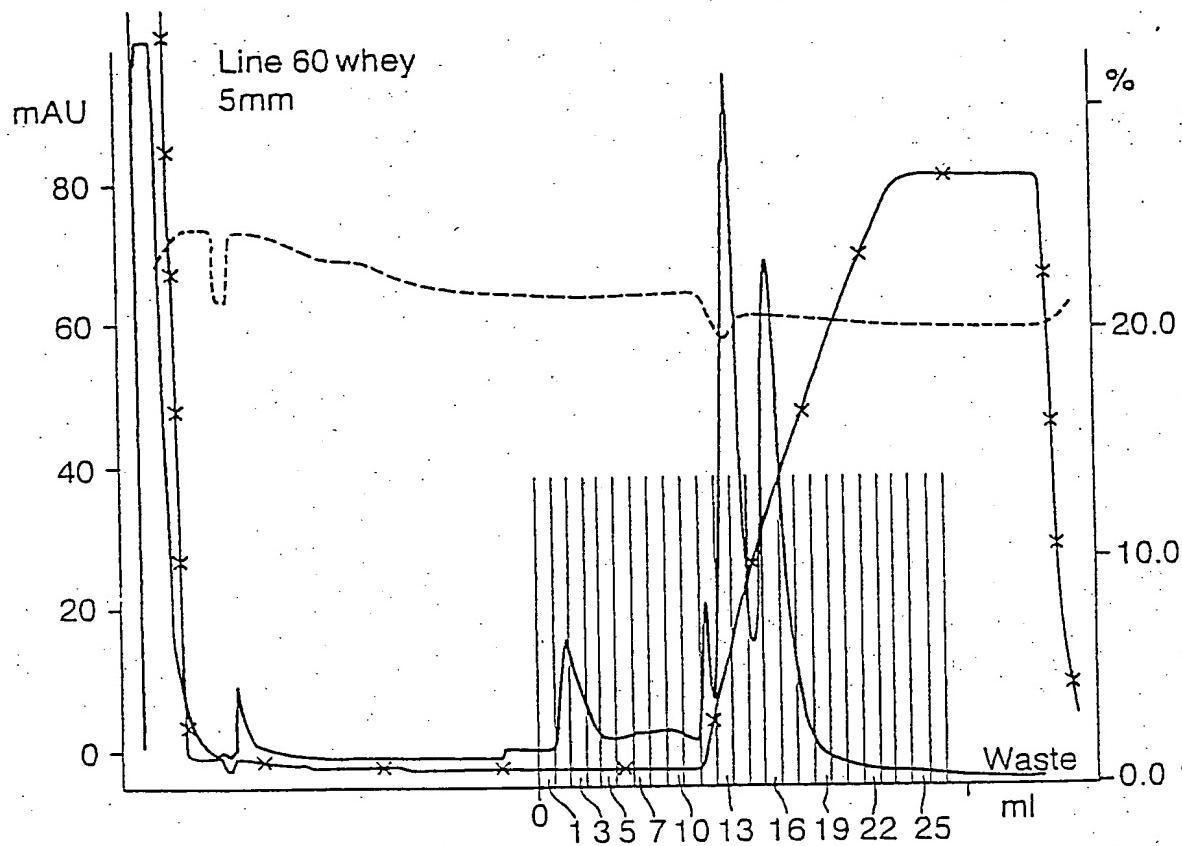


Fig. 15.

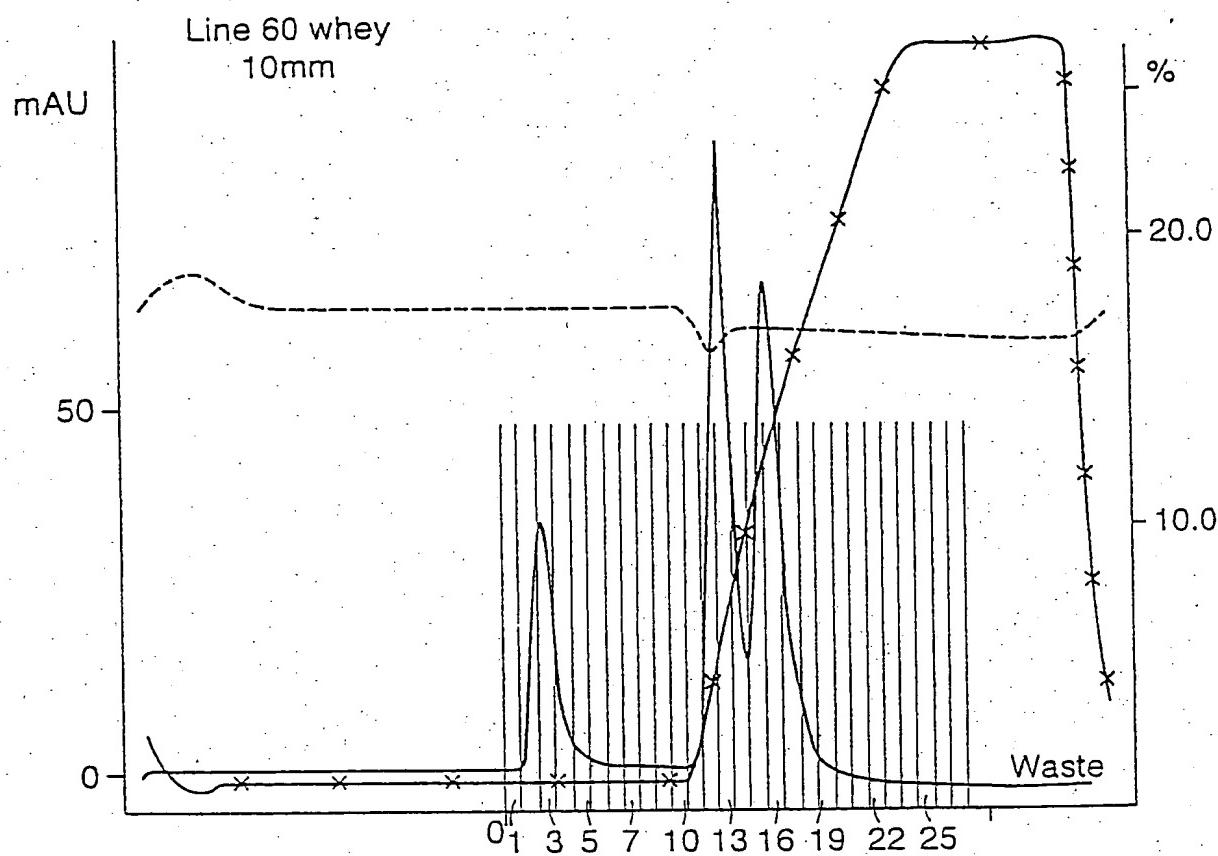
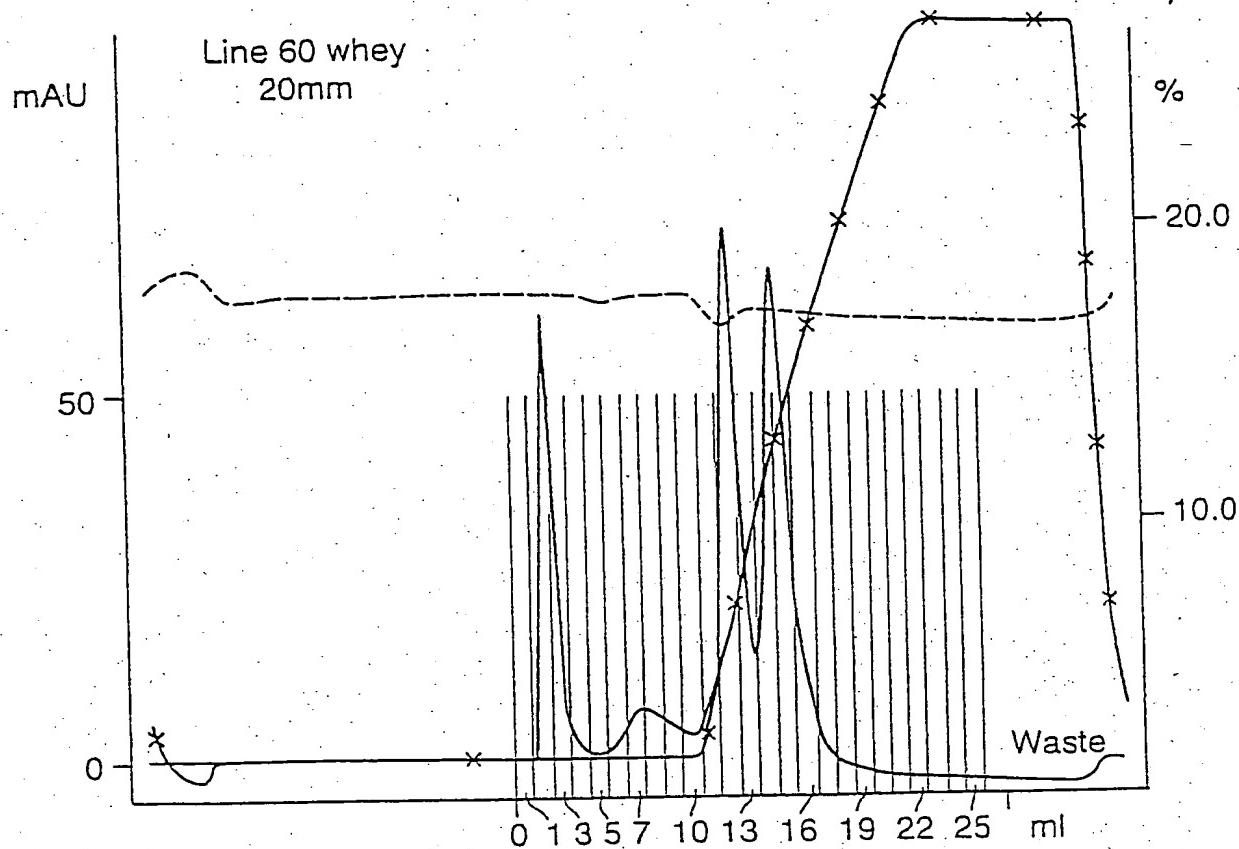


Fig. 16.



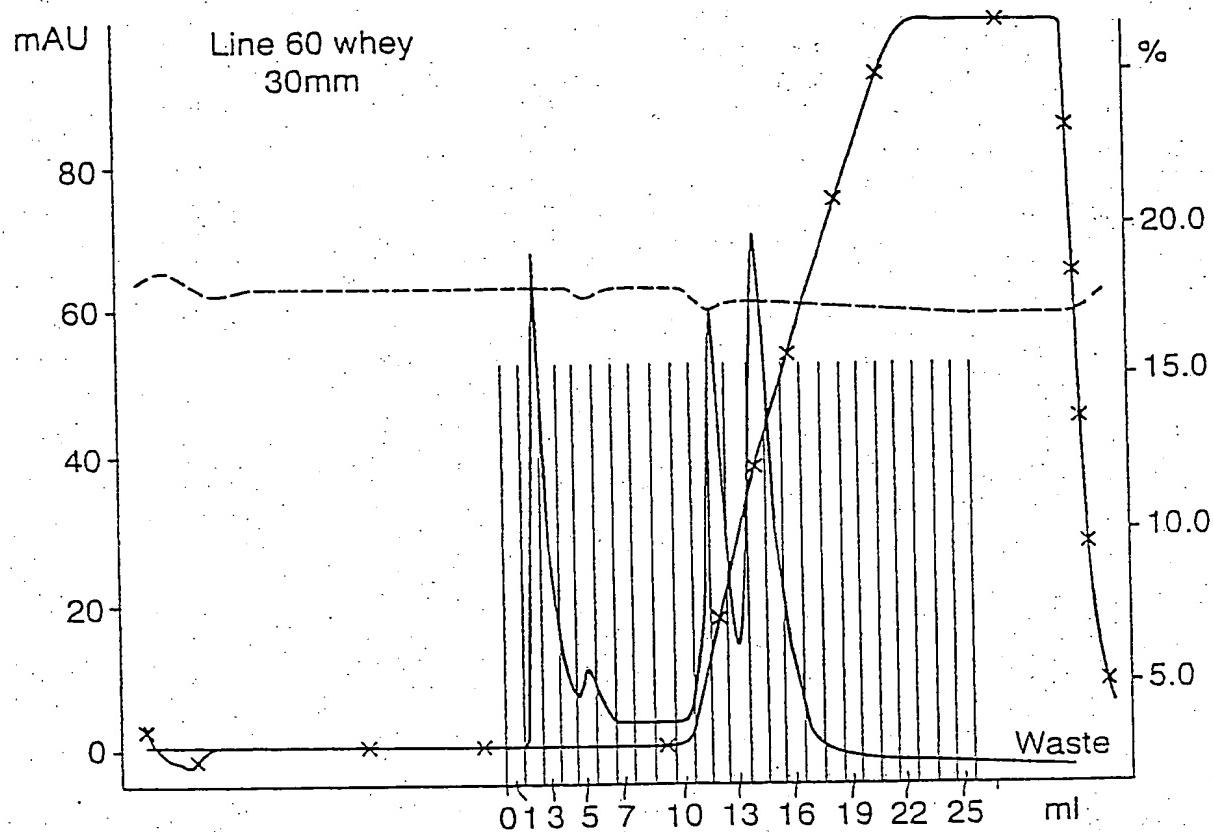
— 12099803:12_UV1_280nm

- - - 12099803:12_pH

* * * 12099803:12_Cond%

12099803:12_Fractions

Fig. 17.



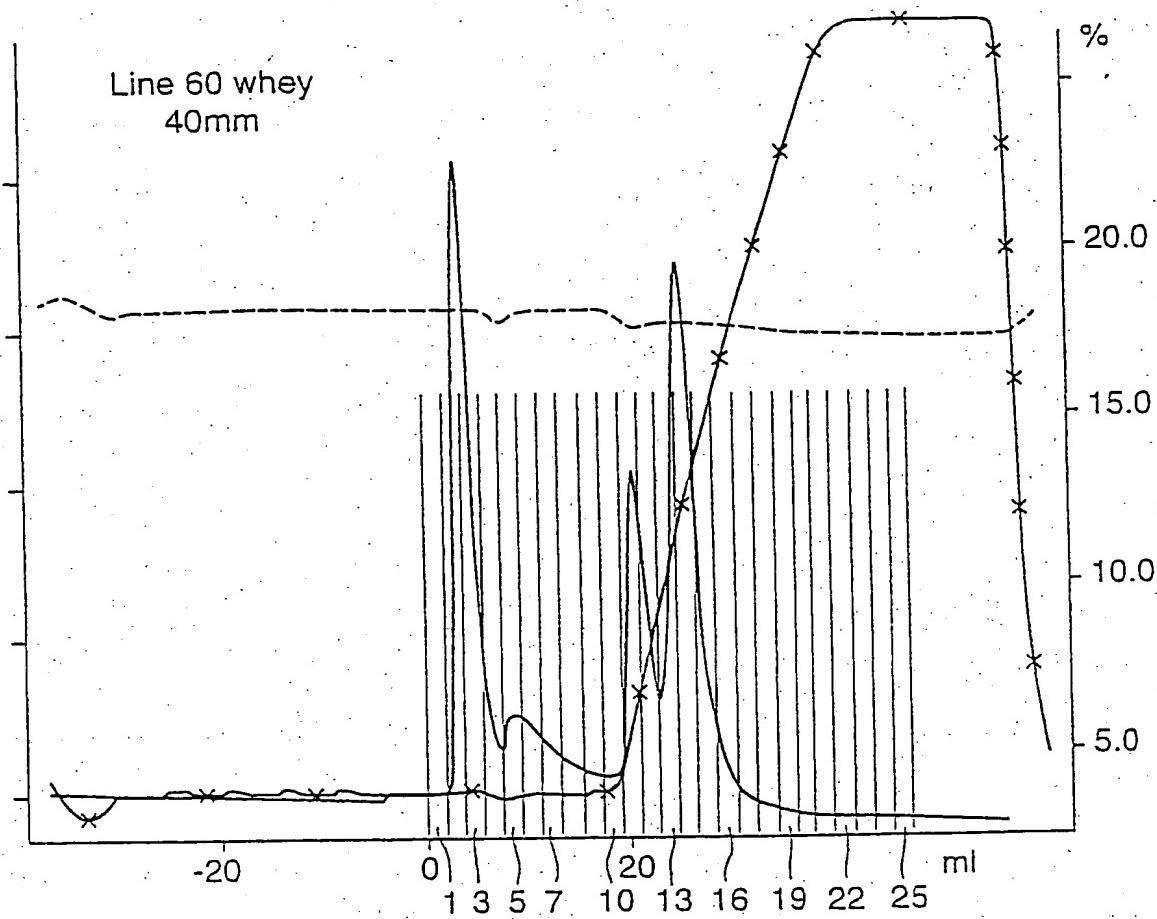
_____ 12099804:13_UV1_280nm

----- 12099804:13_pH

--* 12099804:13_Cond%

12099804:13_Fractions

Fig. 18.



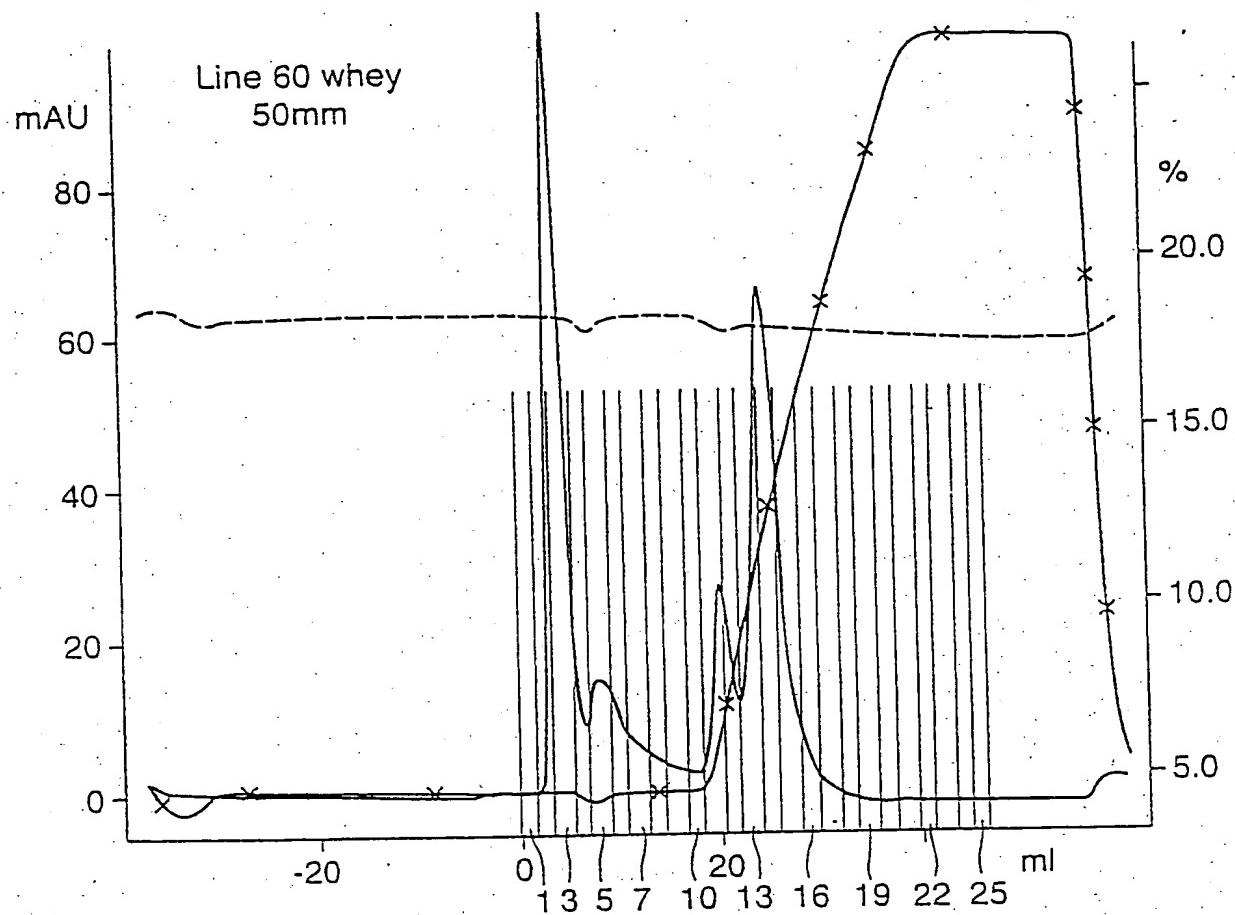
121099805:1_UV1_280nm

121099805:1_pH

121099805:1_Cond%

121099805:1_Fractions

Fig. 19.



——— 121099806:1_UV1_280nm

----- 121099806:1_pH

——*—*—* 121099806:1_Cond%

121099806:1_Fractions

Fig. 20.

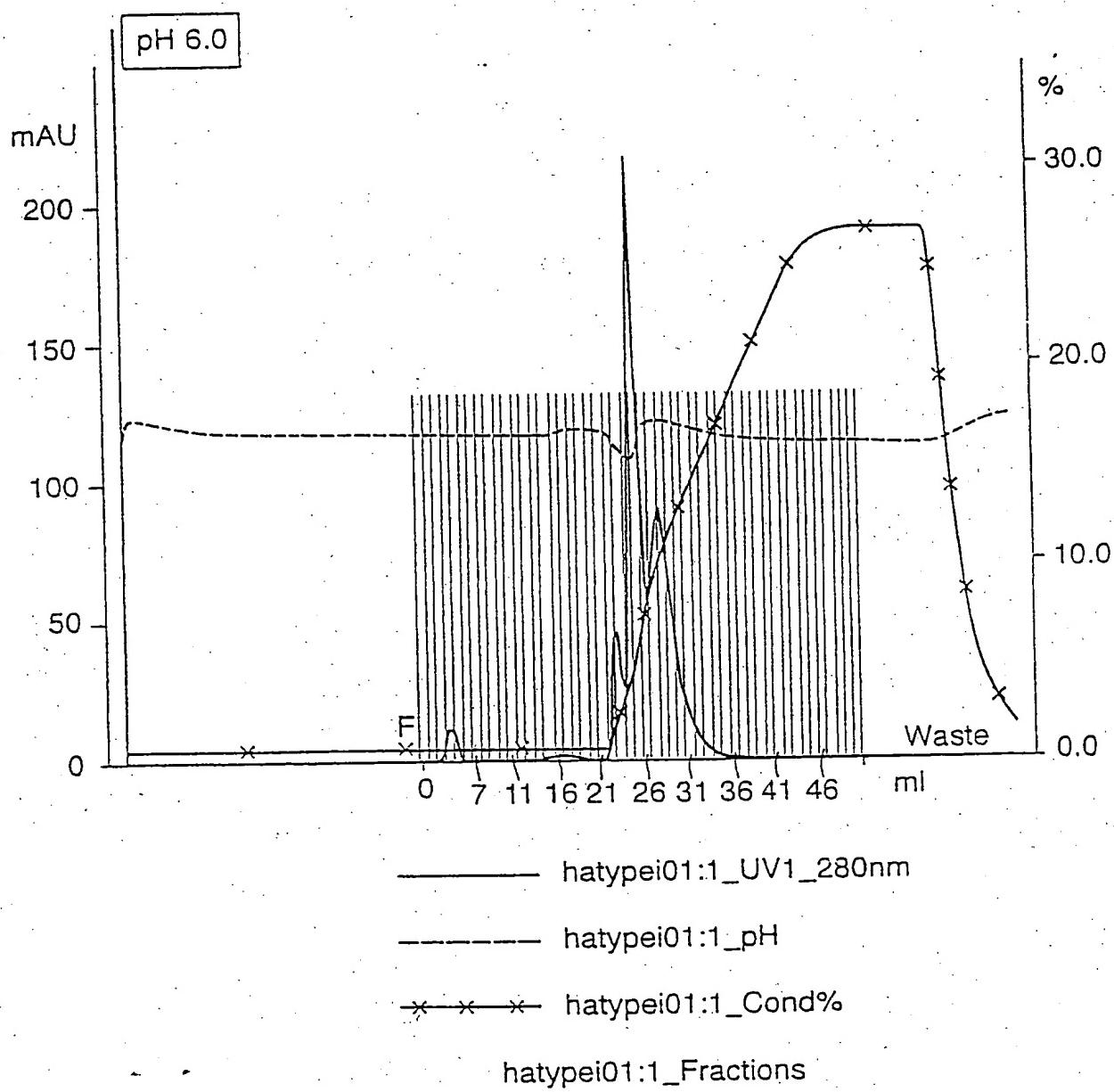


Fig. 21.

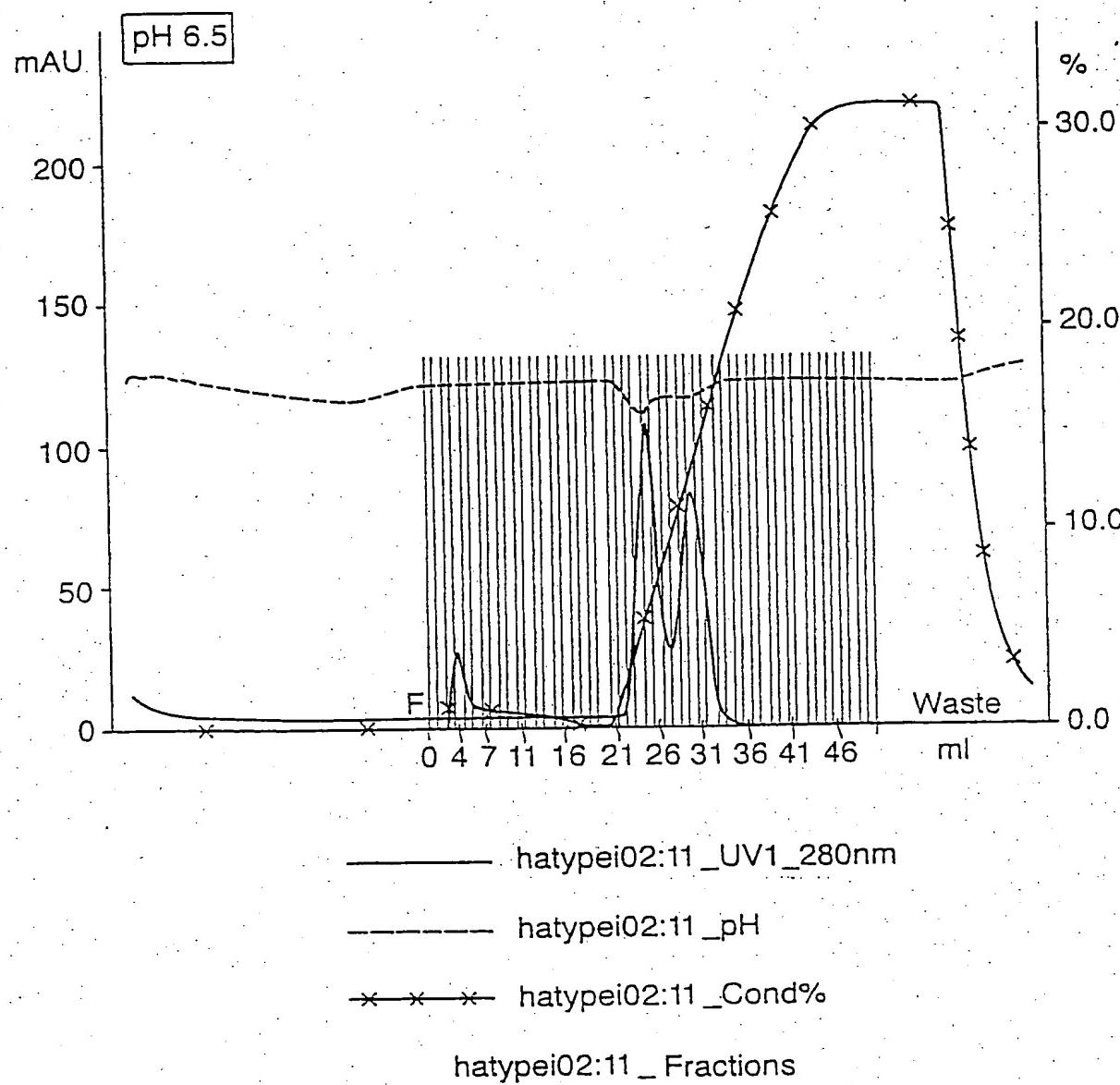
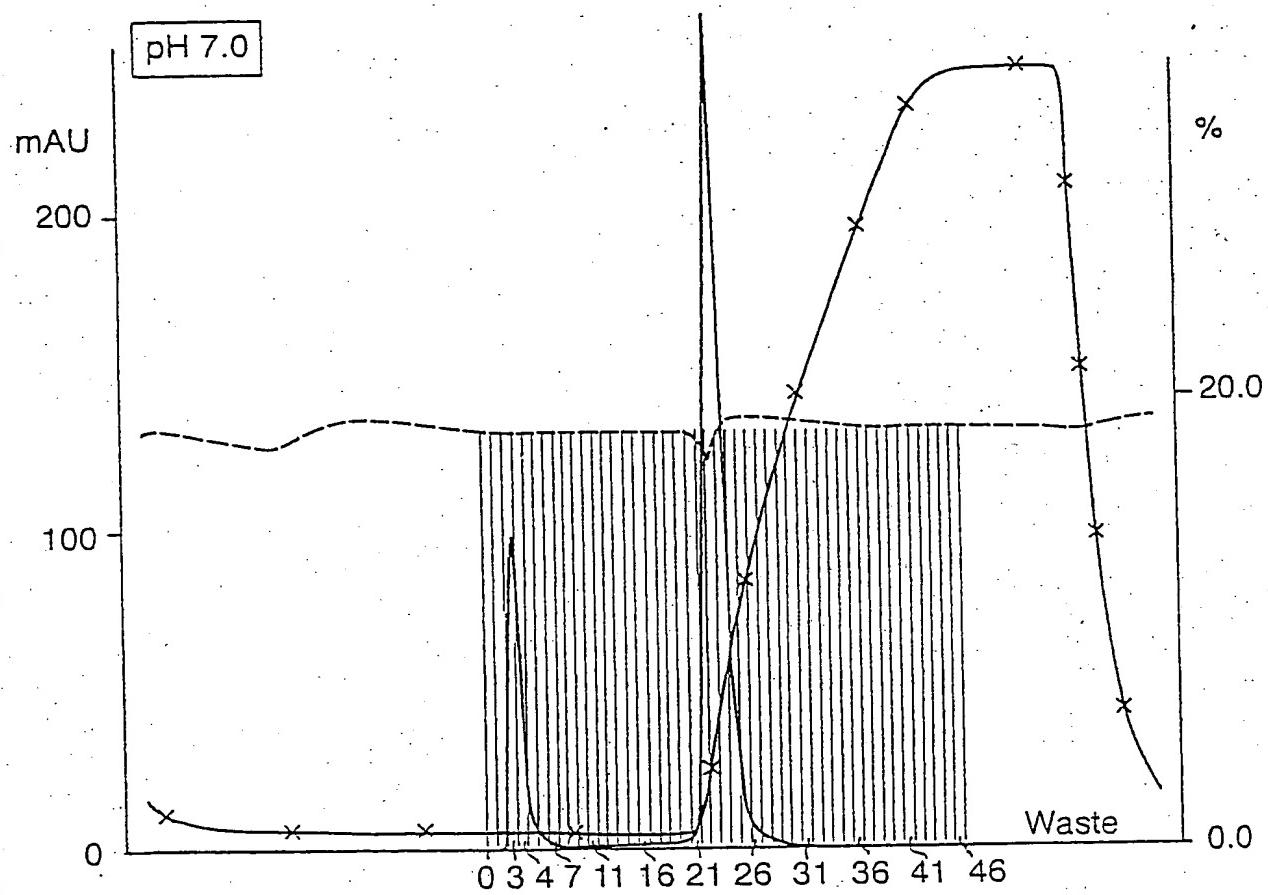


Fig. 22.



_____ hatypei03:12_UV1_280nm

----- hatypei03:12_pH

--* hatypei03:12_Cond%

hatypei03:12_Fractions

Fig. 23.

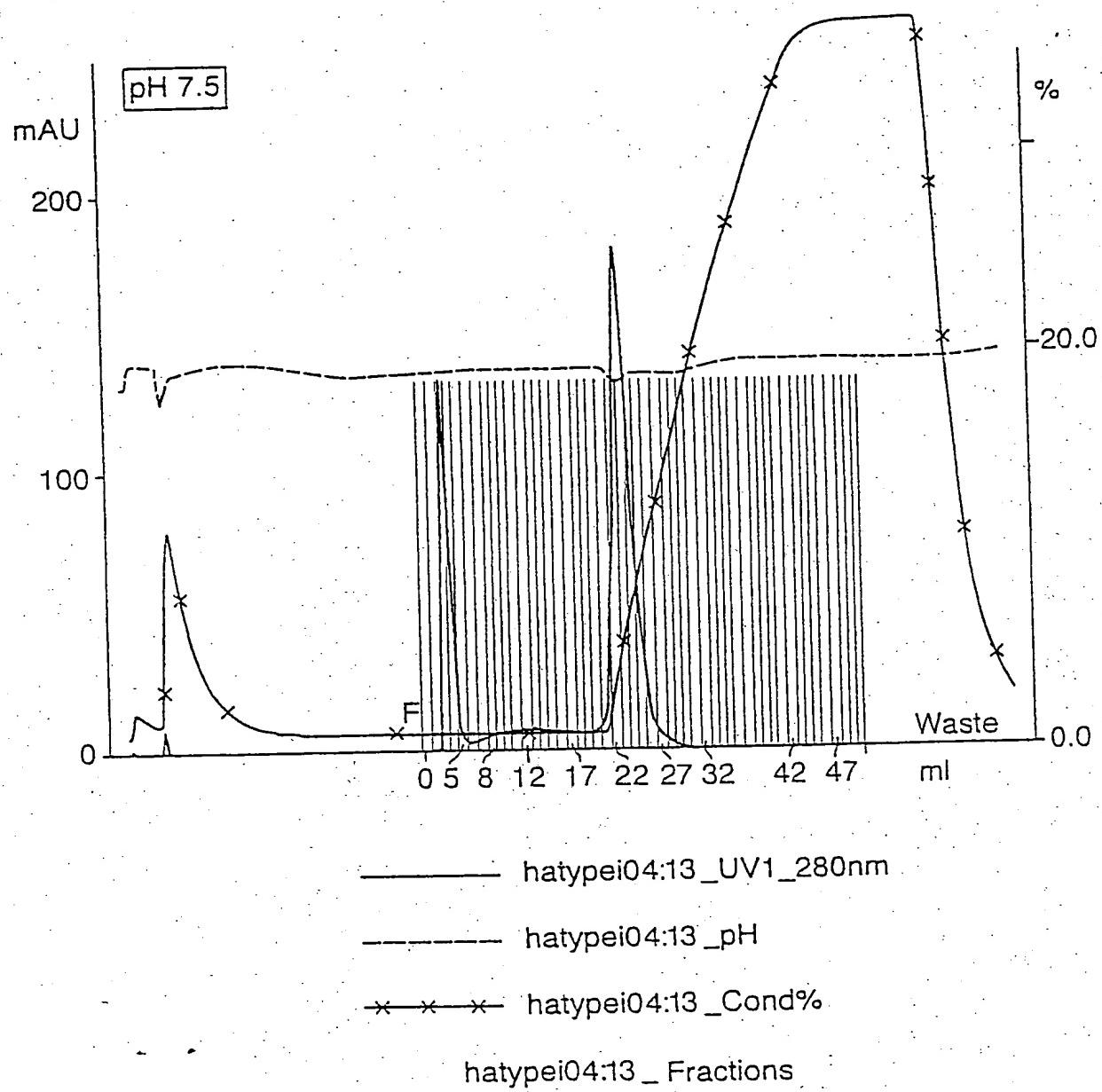


Fig. 24.

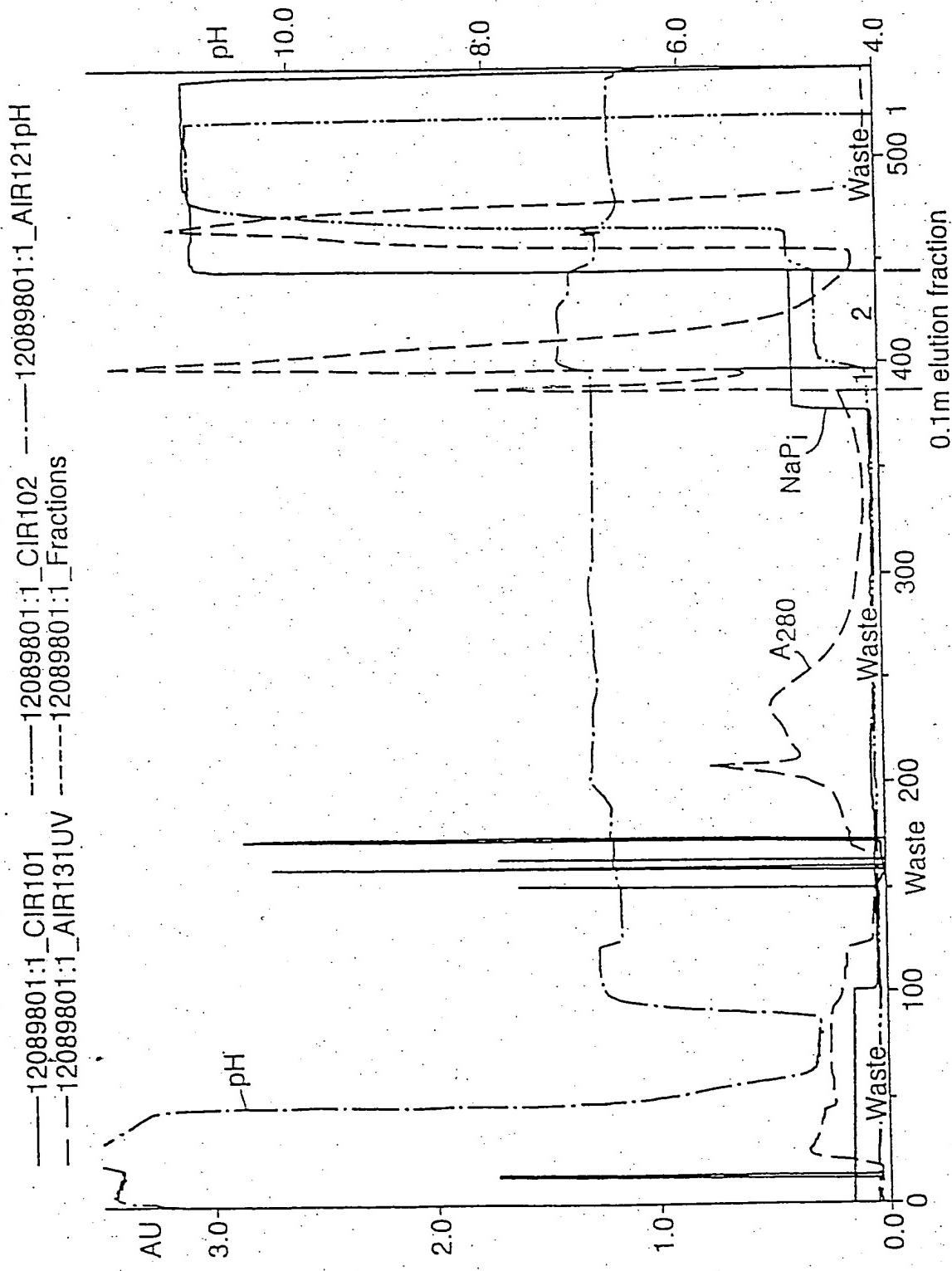


Fig. 25.

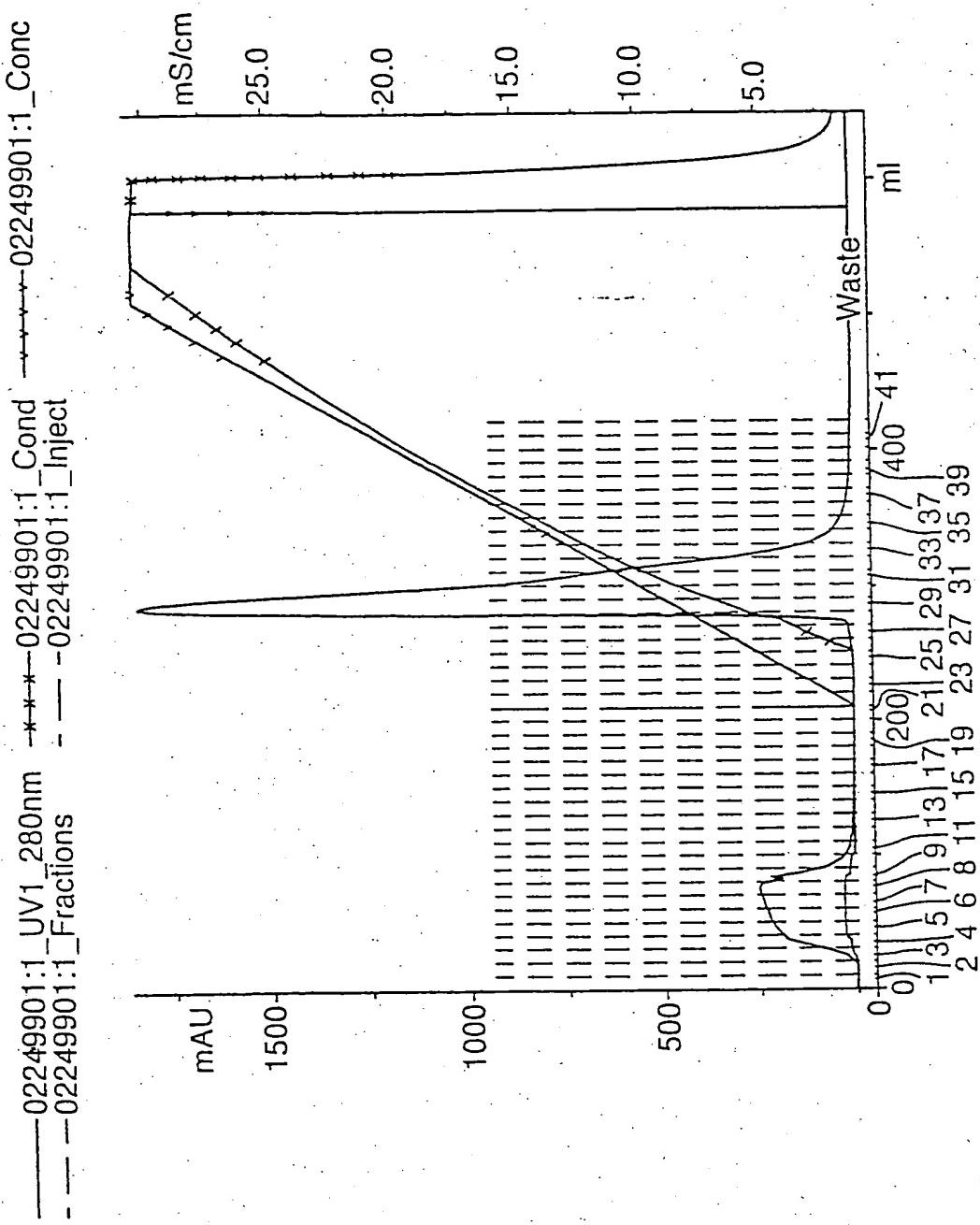


Fig. 26.

XK16/15 80°C
cHT type I 10mM Napi pH 6.5 ; QFF eluate
Run 02249901/02259901/02269901

